



PCT10

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/088,079

DATE: 04/09/2002

TIME: 13:06:45

Input Set : A:\45419_SeqList.txt

Output Set: N:\CRF3\04092002\J088079.raw

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3 <110> APPLICANT: National Research Council of Canada
5 <120> TITLE OF INVENTION: Transgenic manipulation of sn-glycerol-3-phosphate and
6   glycerol production with a feedback defective
7   glycerol-3-phosphate dehydrogenase gene
9 <130> FILE REFERENCE: 45419
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/088,079
C--> 12 <141> CURRENT FILING DATE: 2002-03-21
14 <150> PRIOR APPLICATION NUMBER: US60/155133
15 <151> PRIOR FILING DATE: 1999-09-22
17 <160> NUMBER OF SEQ ID NOS: 6
19 <170> SOFTWARE: PatentIn Ver. 2.1
21 <210> SEQ ID NO: 1
22 <211> LENGTH: 1020
23 <212> TYPE: DNA
24 <213> ORGANISM: Escherichia coli
26 <400> SEQUENCE: 1
27 atgaaccaac gtaatgcttc aatgactgtg atcgggtgccg gctcgtacgg caccgctctt 60
28 gccatcacc tggcaagaaa tggccacgag gttgtcctct ggggccatga ccctgaacat 120
29 atcgcaacgc ttgaacgcga ccgctgtaac gccgcgtttc tccccgatgt gccttttccc 180
30 gatacgtctc atcttgaaag cgatctcgcc actgcgctgg cagccagccg taatattctc 240
31 gtcgtcgtac ccagccatgt ctttggtgaa gtgctgcgcc agattaaacc actgatgcgt 300
32 cctgatgcgc gtctggtgtg ggcgaccaa gggctggaag cggaaaccgg acgtctgtta 360
33 caggacgtgg cgcgtgaggc cttaggcgat caaattccgc tggcggttat ctctggccca 420
34 acgtttgcga aagaactggc ggcaggttta ccgacagcta ttctgctggc ctcgaccgat 480
35 cagacctttg ccgatgatct ccagcagctg ctgcactgcg gcaaaaagttt ccgcgtttac 540
36 agcaatccgg atttcattgg cgtgcagctt ggcggcgcgg tgaaaaacgt tattgccatt 600
37 ggtgcgggga tgtccgacgg tatcggtttt ggtgcgaatg cgcgtacggc gctgatcacc 660
38 cgtgggctgg ctgaaatgtc gcgtcttggt gcggcgctgg gtgccgaccc tgccaccttt 720
39 atgggcatgg cggggcttgg cgatctggtg cttacctgta ccgaaaacca gtcgcgtaac 780
40 cgccgttttg gcatgatgct cggtcagggc atggatgtac aaagcgcgca ggagaagatt 840
41 ggtcaggtgg tggaaggcta ccgcaatacg aaagaagtcc gcgaactggc gcatcgcttc 900
42 ggcgttgaaa tgccaataac cgaggaaatt tatcaagtat tatattgcgg aaaaaacgcg 960
43 cgcgaggcag cattgacttt actaggtcgt gcacgcaagg acgagcgcag cagccactaa 1020
45 <210> SEQ ID NO: 2
46 <211> LENGTH: 339
47 <212> TYPE: PRT
48 <213> ORGANISM: Escherichia coli
50 <400> SEQUENCE: 2
51 Met Asn Gln Arg Asn Ala Ser Met Thr Val Ile Gly Ala Gly Ser Tyr
52   1               5               10               15
54 Gly Thr Ala Leu Ala Ile Thr Leu Ala Arg Asn Gly His Glu Val Val
55           20           25           30
57 Leu Trp Gly His Asp Pro Glu His Ile Ala Thr Leu Glu Arg Asp Arg

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58          35          40          45
60 Cys Asn Ala Ala Phe Leu Pro Asp Val Pro Phe Pro Asp Thr Leu His
61          50          55          60
63 Leu Glu Ser Asp Leu Ala Thr Ala Leu Ala Ala Ser Arg Asn Ile Leu
64 65          70          75          80
66 Val Val Val Pro Ser His Val Phe Gly Glu Val Leu Arg Gln Ile Lys
67          85          90          95
69 Pro Leu Met Arg Pro Asp Ala Arg Leu Val Trp Ala Thr Lys Gly Leu
70          100          105          110
72 Glu Ala Glu Thr Gly Arg Leu Leu Gln Asp Val Ala Arg Glu Ala Leu
73          115          120          125
75 Gly Asp Gln Ile Pro Leu Ala Val Ile Ser Gly Pro Thr Phe Ala Lys
76          130          135          140
78 Glu Leu Ala Ala Gly Leu Pro Thr Ala Ile Ser Leu Ala Ser Thr Asp
79 145          150          155          160
81 Gln Thr Phe Ala Asp Asp Leu Gln Gln Leu Leu His Cys Gly Lys Ser
82          165          170          175
84 Phe Arg Val Tyr Ser Asn Pro Asp Phe Ile Gly Val Gln Leu Gly Gly
85          180          185          190
87 Ala Val Lys Asn Val Ile Ala Ile Gly Ala Gly Met Ser Asp Gly Ile
88          195          200          205
90 Gly Phe Gly Ala Asn Ala Arg Thr Ala Leu Ile Thr Arg Gly Leu Ala
91          210          215          220
93 Glu Met Ser Arg Leu Gly Ala Ala Leu Gly Ala Asp Pro Ala Thr Phe
94 225          230          235          240
96 Met Gly Met Ala Gly Leu Gly Asp Leu Val Leu Thr Cys Thr Glu Asn
97          245          250          255
99 Gln Ser Arg Asn Arg Arg Phe Gly Met Met Leu Gly Gln Gly Met Asp
100          260          265          270
102 Val Gln Ser Ala Gln Glu Lys Ile Gly Gln Val Val Glu Gly Tyr Arg
103          275          280          285
105 Asn Thr Lys Glu Val Arg Glu Leu Ala His Arg Phe Gly Val Glu Met
106          290          295          300
108 Pro Ile Thr Glu Glu Ile Tyr Gln Val Leu Tyr Cys Gly Lys Asn Ala
109 305          310          315          320
111 Arg Glu Ala Ala Leu Thr Leu Leu Gly Arg Ala Arg Lys Asp Glu Arg
112          325          330          335
114 Ser Ser His
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119 <211> LENGTH: 18
120 <212> TYPE: DNA
121 <213> ORGANISM: Artificial Sequence
123 <220> FEATURE:
124 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
125      GPSA3
127 <400> SEQUENCE: 3
128 ttagtggtg ctgcgctc
130 <210> SEQ ID NO: 4
131 <211> LENGTH: 18

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132 <212> TYPE: DNA
133 <213> ORGANISM: Artificial Sequence
135 <220> FEATURE:
136 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
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139 <400> SEQUENCE: 4
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143 <211> LENGTH: 26
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145 <213> ORGANISM: Artificial Sequence
147 <220> FEATURE:
148 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR pimer
149     GPSA31
151 <400> SEQUENCE: 5
152 gagagctctt agtggctgct gcgctc                                26
154 <210> SEQ ID NO: 6
155 <211> LENGTH: 30
156 <212> TYPE: DNA
157 <213> ORGANISM: Artificial Sequence
159 <220> FEATURE:
160 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
161     GPSA51
163 <400> SEQUENCE: 6
164 gaagaaggat ccaacaatga accaacgtaa                                30

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VERIFICATION SUMMARY

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Input Set : A:\45419_SeqList.txt

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L:11 M:270 C: Current Application Number differs, Replaced Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date